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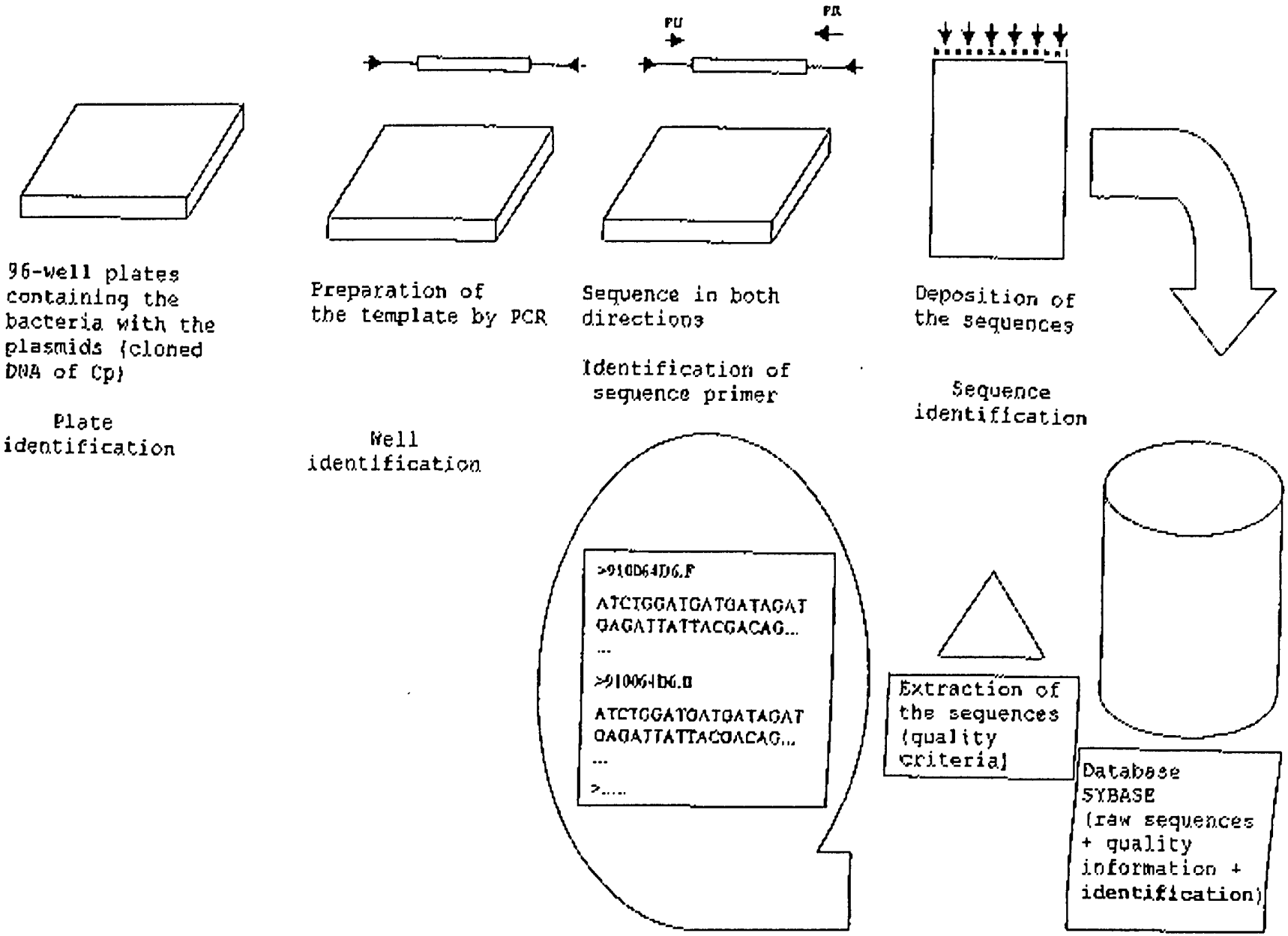

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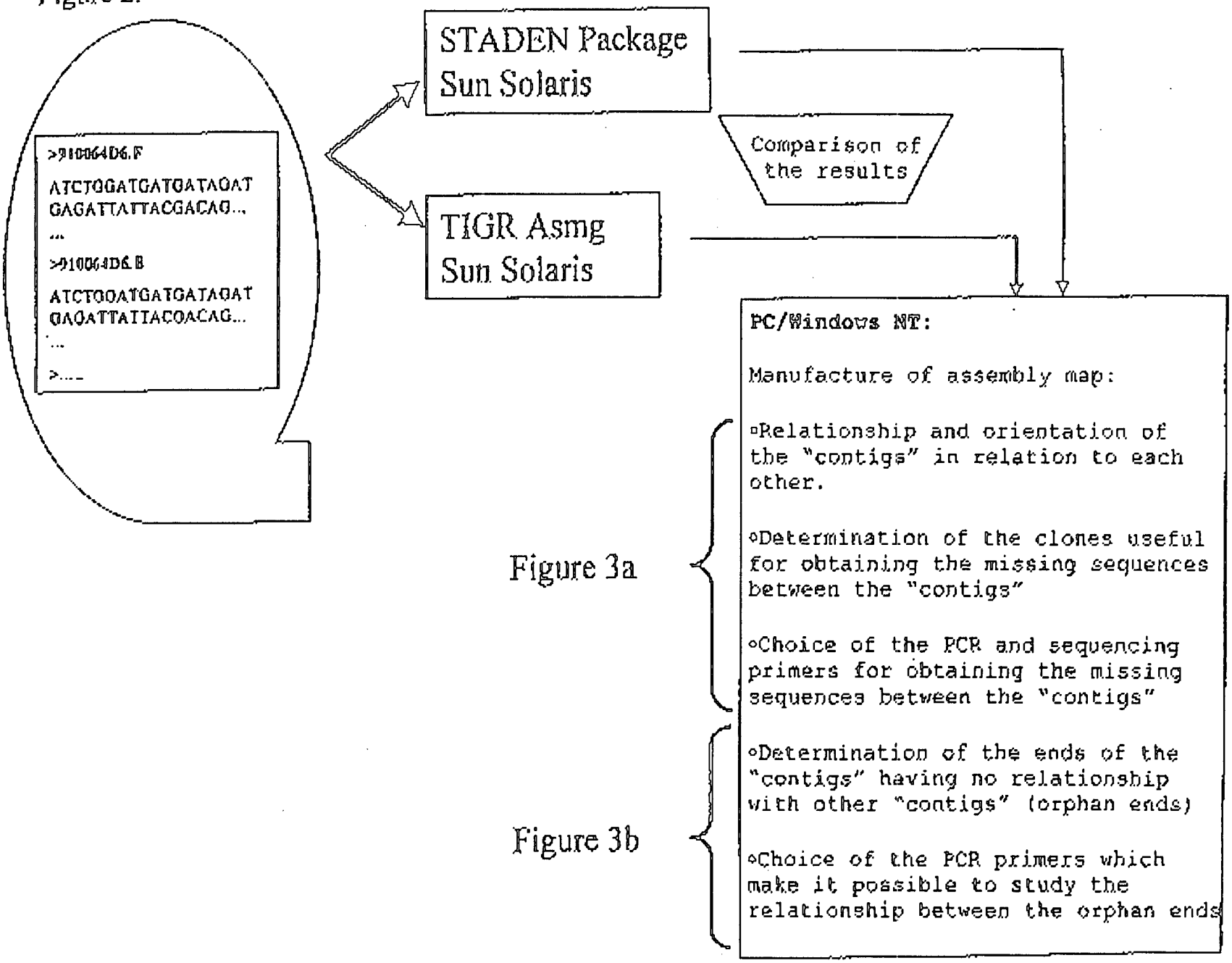

Figure 3a

Figure 3b

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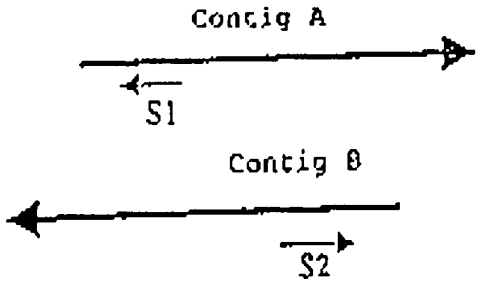
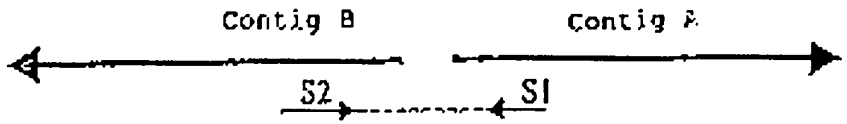
FIGURE 3A



Statistical determination of the sequences:

1- Belonging to the same clone

2- Situated on two different contigs

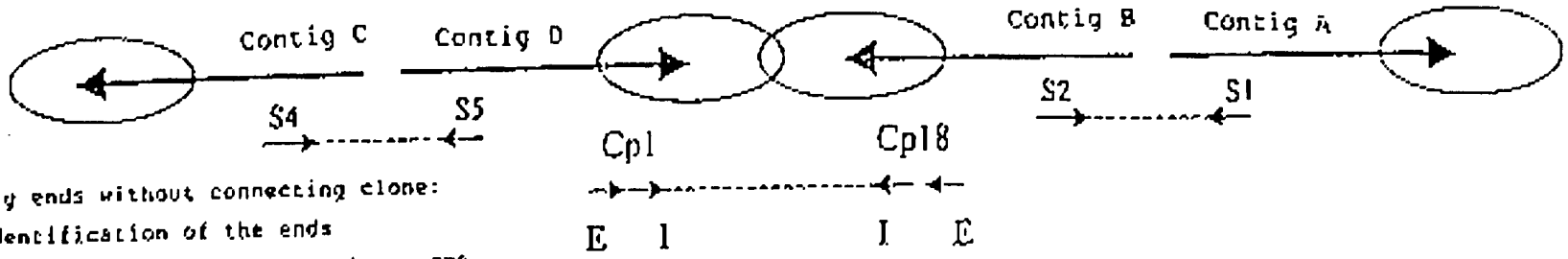


Distribution and position of the sequences on the contigs:

1- Define the arrangement of the contigs in relation to each other

2- Define the PCR primers which make it possible to fill the sequence

FIGURE 3B



Contig ends without connecting clone:

1- Identification of the ends

2- Determination of outer and inner PCR primers for studying the relationships between the contigs

E: outer primers

I: inner primers

SUBSTITUTE SHEET (RULE 26)

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